



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/737,476A

Source: O1PE

Date Processed by STIC: 6/20/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 09/237,476A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 Wrapped Nucleic acids The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

3 Misaligned Amino Acid Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5 Variable Length Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 PatentIn 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7 Skipped Sequences (OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8 Skipped Sequences (NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

9 Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10 Invalid <213> Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11 Use of <220> Sequence(s) SS missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

12 PatentIn 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/737,476A

DATE: 06/20/2001
TIME: 14:38:23

Input Set : A:\09737476.txt
Output Set: N:\CRF3\06202001\I737476A.raw

Does Not Comply
Corrected Diskette Needed

6

3 <110> APPLICANT: Leo G.J. FRENKEN
4 Cornelis P.E. van der LOGT
5 Vin-Miin TEH
6 Martine E. VERHOEYEN
7 Joy E. WILKINSON
8 Stephen A. JOBLING
10 <120> TITLE OF INVENTION: Production of Antibodies
12 <130> FILE REFERENCE: 060113/0275850 - T7060C
14 <140> CURRENT APPLICATION NUMBER: US 09/737,476A
16 <141> CURRENT FILING DATE: 2000-12-18
18 <150> PRIOR APPLICATION NUMBER: EP 99310188.0
20 <151> PRIOR FILING DATE: 1999-12-17
22 <160> NUMBER OF SEQ ID NOS: 67
23 <170> SOFTWARE: MS Word
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 440
27 <212> TYPE: DNA
C--> 28 <213> ORGANISM: Artificial
30 <220> FEATURE:
31 <223> OTHER INFORMATION: VHH with peptide linker
33 <220> FEATURE:
34 <221> NAME/KEY: CDS
35 <222> LOCATION: (1)..(417)
37 <400> SEQUENCE: 1
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40 Gln Val Gln Leu Gln Glu Ser Gly Gly Leu Val Gln Ala Gly Gly
41 1 5 10 15
43. tct ctg aga ctc tcc tgt gca gcc tcg gga cgc gcc acc agt ggt cat 96
44 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Arg Ala Thr Ser Gly His
45 20 25 30
47 ggt cac tat ggt atg ggc tgg ttc cgc cag gtt cca ggg aag gag cgt 144
48 Gly His Tyr Gly Met Gly Trp Phe Arg Gln Val Pro Gly Lys Glu Arg
49 35 40 45
51 gag ttt gtc gca gct att agg tgg agt ggt aaa gag aca tgg tat aaa 192
52 Glu Phe Val Ala Ala Ile Arg Trp Ser Gly Lys Glu Thr Trp Tyr Lys
53 50 55 60
55 gac tcc gtg aag ggc cga ttc acc atc tcc aga gat aac gcc aag act 240
56 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Thr
57 65 70 75 80
59 acg gtt tat ctg caa atg aac agc ctg aaa cct gaa gat acg gcc gtt 288
60 Thr Val Tyr Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val
61 85 90 95
63 tat tat tgt gcc gct cga ccg gtc cgc gtg gat gat att tcc ctg ccg 336
64 Tyr Tyr Cys Ala Ala Arg Pro Val Arg Val Asp Asp Ile Ser Leu Pro
65 100 105 110
67 gtt ggg ttt gac tac tgg ggc cag ggg acc cag gtc acc gtc tcc tca 384
68 Val Gly Phe Asp Tyr Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser

RAW SEQUENCE LISTING
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Input Set : A:\09737476.txt
Output Set: N:\CRF3\06202001\I737476A.raw

69 115 120 125
71 gaa caa aaa ctc atc tca gaa gag gat ctg aat taataaggc taagctcgaa 437
72 Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn
73 130 135
75 ttc 440
78 <210> SEQ ID NO: 2
79 <211> LENGTH: 139
80 <212> TYPE: PRT
C--> 81 <213> ORGANISM: Artificial
83 <220> FEATURE:
84 <223> OTHER INFORMATION: VHH with peptide linker
86 <400> SEQUENCE: 2
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89 1 5 10 15
91 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Arg Ala Thr Ser Gly His
92 20 25 30
94 Gly His Tyr Gly Met Gly Trp Phe Arg Gln Val Pro Gly Lys Glu Arg
95 35 40 45
97 Glu Phe Val Ala Ala Ile Arg Trp Ser Gly Lys Glu Thr Trp Tyr Lys
98 50 55 60
100 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Thr
101 65 70 75 80
103 Thr Val Tyr Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val
104 85 90 95
106 Tyr Tyr Cys Ala Ala Arg Pro Val Arg Val Asp Asp Ile Ser Leu Pro
107 100 105 110
109 Val Gly Phe Asp Tyr Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser
110 115 120 125
112 Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn
113 130 135
116 <210> SEQ ID NO: 3
117 <211> LENGTH: 11
118 <212> TYPE: PRT
C--> 119 <213> ORGANISM: Artificial
121 <220> FEATURE:
122 <223> OTHER INFORMATION: myc linker
124 <400> SEQUENCE: 3
126 Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn
127 1 5 10
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131 <211> LENGTH: 471
132 <212> TYPE: DNA
C--> 133 <213> ORGANISM: Artificial
135 <220> FEATURE:
136 <223> OTHER INFORMATION: VHH with linker
138 <220> FEATURE:
139 <221> NAME/KEY: CDS
140 <222> LOCATION: (1)..(459)
142 <400> SEQUENCE: 4

RAW SEQUENCE LISTING
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DATE: 06/20/2001
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Output Set: N:\CRF3\06202001\I737476A.raw

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145	Gln	Val	Gln	Leu	Gln	Gln	Ser	Gly	Gly	Gly	Leu	Val	Gln	Ala	Gly	Gly	
146	1			5				10						15			
148	tct	ctg	aga	ctc	tcc	tgt	gta	gct	tct	gaa	agc	agc	ttc	agc	agc	aat	96
149	Ser	Leu	Arg	Leu	Ser	Cys	Val	Ala	Ser	Glu	Ser	Ser	Phe	Ser	Asn	Asn	
150				20				25					30				
152	cac	atg	ggc	tgg	tac	cgc	cgg	gct	cca	ggg	aac	cag	cgc	gag	ctg	gtc	144
153	His	Met	Gly	Trp	Tyr	Arg	Arg	Ala	Pro	Gly	Asn	Gln	Arg	Glu	Leu	Val	
154		35					40					45					
156	gca	act	att	agt	cct	ggt	ggt	agc	aca	cac	tat	gta	gac	tcc	gtg	aag	192
157	Ala	Thr	Ile	Ser	Pro	Gly	Gly	Ser	Thr	His	Tyr	Val	Asp	Ser	Val	Lys	
158		50					55					60					
160	ggc	cga	ttc	acc	atc	tcc	cga	gac	aac	gcc	aag	aac	aca	gtg	tat	cta	240
161	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ala	Lys	Asn	Thr	Val	Tyr	Leu	
162	65				70					75			80				
164	caa	atg	gac	agc	ctg	aaa	cca	gag	gac	acg	gcc	gtc	tat	tac	tgt	gct	288
165	Gln	Met	Asp	Ser	Leu	Lys	Pro	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys	Ala	
166		85					90				95						
168	gcc	aag	ggg	agg	ggg	ctg	cag	gct	atg	cag	tac	tgg	ggc	cag	ggg	acc	336
169	Ala	Lys	Gly	Arg	Gly	Leu	Gln	Ala	Met	Gln	Tyr	Trp	Gly	Gln	Gly	Thr	
170		100					105				110						
172	ctg	gtc	acc	gtc	tcc	tca	gct	cac	cac	agc	gaa	gac	ccc	agc	tcc	gct	384
173	Leu	Val	Thr	Val	Ser	Ser	Ala	His	His	Ser	Glu	Asp	Pro	Ser	Ser	Ala	
174		115					120				125						
176	gcc	gcc	cat	cac	cat	cac	cat	cac	ggg	gcc	gca	caa	aaa	ctc	atc	1432	
177	Ala	Ala	His	His	His	His	His	His	Gly	Ala	Ala	Glu	Gln	Lys	Leu	Ile	
178		130					135				140						
180	tca	gaa	gag	gat	ctg	aat	ggg	gcc	gca	tagtaacaat	tg					471	
181	Ser	Glu	Glu	Asp	Leu	Asn	Gly	Ala	Ala								
182	145				150												
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186	<211> LENGTH: 153																
187	<212> TYPE: PRT																

C--> 188 <213> ORGANISM: Artificial

190 <220> FEATURE:

191 <223> OTHER INFORMATION: VHH with linker

193 <400> SEQUENCE: 5

195	Gln	Val	Gln	Leu	Gln	Gln	Ser	Gly	Gly	Gly	Leu	Val	Gln	Ala	Gly	Gly	
196	1			5				10					15				
198	Ser	Leu	Arg	Leu	Ser	Cys	Val	Ala	Ser	Glu	Ser	Ser	Phe	Ser	Asn	Asn	
199		20					25				30						
201	His	Met	Gly	Trp	Tyr	Arg	Arg	Ala	Pro	Gly	Asn	Gln	Arg	Glu	Leu	Val	
202		35					40				45						
204	Ala	Thr	Ile	Ser	Pro	Gly	Gly	Ser	Thr	His	Tyr	Val	Asp	Ser	Val	Lys	
205		50					55				60						
207	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ala	Lys	Asn	Thr	Val	Tyr	Leu	
208	65				70					75			80				
210	Gln	Met	Asp	Ser	Leu	Lys	Pro	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys	Ala	
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PATENT APPLICATION: US/09/737,476A

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Input Set : A:\09737476.txt
Output Set: N:\CRF3\06202001\I737476A.raw

213 Ala Lys Gly Arg Gly Leu Gln Ala Met Gln Tyr Trp Gly Gln Gly Thr
214 100 105 110
216 Leu Val Thr Val Ser Ser Ala His His Ser Glu Asp Pro Ser Ser Ala
217 115 120 125
219 Ala Ala His His His His Gly Ala Ala Glu Gln Lys Leu Ile
220 130 135 140
222 Ser Glu Glu Asp Leu Asn Gly Ala Ala
223 145 150
226 <210> SEQ ID NO: 6
227 <211> LENGTH: 468
228 <212> TYPE: DNA

C--> 229 <213> ORGANISM: Artificial

231 <220> FEATURE:
232 <223> OTHER INFORMATION: VHH with linker
234 <220> FEATURE:
235 <221> NAME/KEY: CDS
236 <222> LOCATION: (1)..(456)
238 <400> SEQUENCE: 6

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242 1	5									10		15					
244 tct	ctg	aga	ctc	tcc	tgt	gta	gcc	tct	gga	aac	acc	ttc	agt	atc	ata	96	
245 Ser	Leu	Arg	Leu	Ser	Cys	Val	Ala	Ser	Gly	Asn	Thr	Phe	Ser	Ile	Ile		
246	20									25		30					
248 gct	atg	gcc	tgg	tac	cgc	cag	gct	cca	ggg	aag	cag	cgc	gag	gtg	gtc	144	
249 Ala	Met	Ala	Trp	Tyr	Arg	Gln	Ala	Pro	Gly	Lys	Gln	Arg	Glu	Val	Val		
250	35									40		45					
252 gca	agt	att	aat	agt	att	ggc	agc	aca	aat	tat	gca	gac	tcc	gtg	aag	192	
253 Ala	Ser	Ile	Asn	Ser	Ile	Gly	Ser	Thr	Asn	Tyr	Ala	Asp	Ser	Val	Lys		
254	50									55		60					
256 ggg	cga	ttc	acc	atc	tcc	aga	gac	aac	gcc	aag	aac	aca	gtg	tat	ctg	240	
257 Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ala	Lys	Asn	Thr	Val	Tyr	Leu		
258 65										70		75		80			
260 caa	atg	agc	agc	ctg	aaa	cct	gag	gac	acg	gcc	gtc	tat	tac	tgt	gct	288	
261 Gln	Met	Ser	Ser	Leu	Lys	Pro	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys	Ala		
262	85									90		95					
264 gcc	ggt	aat	ttg	ctg	gtt	aag	agg	cct	tac	tgg	gcc	cag	ggg	acc	ctg	336	
265 Ala	Gly	Asn	Leu	Leu	Val	Lys	Arg	Pro	Tyr	Trp	Gly	Gln	Gly	Thr	Leu		
266	100									105		110					
268 gtc	acc	gtc	tcc	tca	gaa	ccc	aag	aca	cca	aaa	cca	caa	cca	gcg	gcc	384	
269 Val	Thr	Val	Ser	Ser	Glu	Pro	Lys	Thr	Pro	Lys	Pro	Gln	Pro	Ala	Ala		
270	115									120		125					
272 gcc	cat	cac	cat	cac	cat	cac	ggg	gcc	gca	gaa	caa	aaa	ctc	atc	tca	432	
273 Ala	His	His	His	His	His	His	Gly	Ala	Ala	Glu	Gln	Lys	Leu	Ile	Ser		
274	130									135		140					
276 gaa	gag	gat	ctg	aat	ggg	gcc	gca	tagtaacaat	tg							468	
277 Glu	Glu	Asp	Leu	Asn	Gly	Ala	Ala										
278 145										150							
281 <210>	SEQ	ID	NO:	7													

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/737,476A
 DATE: 06/20/2001
 TIME: 14:38:23
 Input Set : A:\09737476.txt
 Output Set: N:\CRF3\06202001\I737476A.raw

282 <211> LENGTH: 152
 283 <212> TYPE: PRT
C--> 284 <213> ORGANISM: Artificial
 286 <220> FEATURE:
 287 <223> OTHER INFORMATION: VHH with linker
 289 <400> SEQUENCE: 7
 291 Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly
 292 1 5 10 15
 294 Ser Leu Arg Leu Ser Cys Val Ala Ser Gly Asn Thr Phe Ser Ile Ile
 295 20 25 30
 297 Ala Met Ala Trp Tyr Arg Gln Ala Pro Gly Lys Gln Arg Glu Val Val
 298 35 40 45
 300 Ala Ser Ile Asn Ser Ile Gly Ser Thr Asn Tyr Ala Asp Ser Val Lys
 301 50 55 60
 303 Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Val Tyr Leu
 304 65 70 75 80
 306 Gln Met Ser Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys Ala
 307 85 90 95
 309 Ala Gly Asn Leu Leu Val Lys Arg Pro Tyr Trp Gly Gln Gly Thr Leu
 310 100 105 110
 312 Val Thr Val Ser Ser Glu Pro Lys Thr Pro Lys Pro Gln Pro Ala Ala
 313 115 120 125
 315 Ala His His His His His Gly Ala Ala Glu Gln Lys Leu Ile Ser
 316 130 135 140
 318 Glu Glu Asp Leu Asn Gly Ala Ala
 319 145 150
 322 <210> SEQ ID NO: 8
 323 <211> LENGTH: 462
 324 <212> TYPE: DNA
C--> 325 <213> ORGANISM: Artificial
 327 <220> FEATURE:
 328 <223> OTHER INFORMATION: VHH with linker
 330 <220> FEATURE:
 331 <221> NAME/KEY: CDS
 332 <222> LOCATION: (1)..(450)
 334 <400> SEQUENCE: 8
 336 acc atg gcc cag gtg aaa ctg cag cag tct ggg gga ttg gtg cag 48
 337 Thr Met Ala Gln Val Lys Leu Gln Gln Ser Gly Gly Gly Leu Val Gln
 338 1 5 10 15
 340 gct ggg ggc cct ctg agg ctc tcc tgt gca gcc tct gga cgc acc ttc 96
 341 Ala Gly Gly Pro Leu Arg Leu Ser Cys Ala Ala Ser Gly Arg Thr Phe
 342 20 25 30
 344 agt aac tat gcc gtg ggc tgg ttc cgc cag gct cca ggg aag gag cgt 144
 345 Ser Asn Tyr Ala Val Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg
 346 35 40 45
 348 gag ttt gtc gct gct att agc cgt gat ggt ggg cgc aca tac tat gcg 192
 349 Glu Phe Val Ala Ala Ile Ser Arg Asp Gly Gly Arg Thr Tyr Tyr Ala
 350 50 55 60
 352 gac tcc gtg aag ggc cga ttc gcc gtc tcc aga gac tac gcc gag aac 240

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<210> 55
<211> 377
<212> PRT
<213> Artificial
<400> 55

Met Glu Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly

see item 11 on Enc Summary Sheet

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/737,476A

DATE: 06/20/2001
TIME: 14:38:24

Input Set : A:\09737476.txt
Output Set: N:\CRF3\06202001\I737476A.raw

L:28 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:81 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:119 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:133 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:188 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:229 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:284 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:325 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:380 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:421 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:476 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
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L:765 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:29
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L:791 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:31
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L:873 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:35
L:899 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:36
L:912 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:37
L:925 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:38
L:938 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:39
L:951 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:40
L:967 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:41
L:980 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:42
L:993 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:43
L:1006 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:44
L:1019 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:45
L:1032 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:46
L:1045 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:47
L:1058 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:48

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/737,476A

DATE: 06/20/2001
TIME: 14:38:24

Input Set : A:\09737476.txt
Output Set: N:\CRF3\06202001\I737476A.raw

L:1071 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:49
L:1084 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:50
L:1249 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1249 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: